



Year: 2015

Serotonin transporter and tryptophan hydroxylase gene variations mediate working memory deficits of cocaine users

Havranek, M M ; Vonmoos, Matthias ; Müller, C P ; Büetiger, J R ; Tasiudi, Eve ; Hulka, Lea M ;
Preller, Katrin H ; Mössner, R ; Grünblatt, Edna ; Seifritz, Erich ; Quednow, Boris B

Abstract: Cocaine users consistently develop working memory (WM) impairments but the mediating molecular mechanisms are unknown so far. Recent evidence suggests that the serotonin (5-HT) system is altered by chronic cocaine use, while also being involved in WM processing. Thus, we investigated the effects of genetic variations impacting 5-HT activity and of peripheral 5-HT transporter (5-HTT) mRNA expression on WM performance in cocaine users and stimulant naive controls. Two hundred twenty participants (126 cocaine users, 94 controls) were assessed with visuospatial, spatial, and verbal WM tasks, genotyped for the length polymorphism in the promoter region of the 5-HTT (5-HTTLPR), the variable number of tandem repeats in the second intron of the 5-HTT (VNTR In2), two single-nucleotide polymorphisms (rs4570625 and rs1386497) in the tryptophan hydroxylase-2 (TPH2) gene and quantified for peripheral 5-HTT mRNA expression in whole-blood samples. Several significant gene × environment interactions between 5-HT genotypes and cocaine use on WM emerged: in cocaine users, the long/long (5-HTTLPR), 9+10/9+10 (VNTR In2) and C/C (TPH2 rs1386497) genotypes were risk alleles for WM impairments, whereas in healthy controls these polymorphisms were associated with improved WM performance. Analogously, high 5-HTT mRNA levels were associated with worse executive WM performance in cocaine users but with increased performance in controls. These gene × environment interactions suggest that the 5-HT system has an important role in the development of cognitive deficits in chronic cocaine users. Hence, pharmacological compounds targeting 5-HT neurotransmission might be promising for the treatment of cognitive deficits in cocaine dependence.

DOI: <https://doi.org/10.1038/npp.2015.146>

Posted at the Zurich Open Repository and Archive, University of Zurich

ZORA URL: <https://doi.org/10.5167/uzh-117824>

Journal Article

Accepted Version

Originally published at:

Havranek, M M; Vonmoos, Matthias; Müller, C P; Büetiger, J R; Tasiudi, Eve; Hulka, Lea M; Preller, Katrin H; Mössner, R; Grünblatt, Edna; Seifritz, Erich; Quednow, Boris B (2015). Serotonin transporter and tryptophan hydroxylase gene variations mediate working memory deficits of cocaine users. *Neuropsychopharmacology*, 40:2929-2937.

DOI: <https://doi.org/10.1038/npp.2015.146>

Serotonin transporter and tryptophan hydroxylase gene variations mediate working memory deficits of cocaine users

Michael M Havranek, MD^a, Matthias Vonmoos, PhD^b, Christian P Müller, PhD^c,
Jessica R Büetiger, BSc^b, Eve Tasiudi, MSc^d, Lea M Hulka, PhD^b, Katrin H Preller, PhD^b,
Rainald Mössner, MD^e, Edna Grünblatt, PhD^{d,f}, Erich Seifritz, MD^{a,f}, Boris B Quednow, PhD^{b,f,*}

^a*Department of Psychiatry, Psychotherapy and Psychosomatics, Psychiatric Hospital of the University of Zurich, Switzerland*

^b*Experimental and Clinical Pharmacopsychology, Department of Psychiatry, Psychotherapy and Psychosomatics, Psychiatric Hospital of the University of Zurich, Switzerland*

^c*Department of Psychiatry and Psychotherapy, Friedrich-Alexander-University Erlangen-Nuremberg, Erlangen, Germany*

^d*University Clinic for Child and Adolescent Psychiatry, University of Zurich, Switzerland*

^e*Department of Psychiatry and Psychotherapy, University of Tuebingen, Germany*

^f*Neuroscience Center Zurich, University and ETH Zurich, Switzerland*

Abbreviated Running Title

Serotonin genes mediate working memory deficits in cocaine users

Manuscript Category

Original Article

Resubmitted

May 5th, 2015

Manuscript Characteristics

Number of words in the abstract: 238 / 250

Number of words in the article: 3939 / 4000

Number of figures, tables: 2, 2

Supplemental Material: yes

Disclosures and Acknowledgments

Please see at the end of the text.

Financial support

This study was supported by grants from the Swiss National Science Foundation (SNSF; grant No. PP00P1-123516/1 and PP00P1-146326/1) and the Olga Mayenfisch Foundation.

*Corresponding author

Boris B. Quednow, PhD

Experimental and Clinical Pharmacopsychology

Psychiatric Hospital of the University of Zurich

Lenggstrasse 31, CH-8032 Zurich, Switzerland

Tel.: 0041-44-384-2777 / Fax: 0041-44-384-3396

E-Mail: quedenow@bli.uzh.ch

Abstract

Cocaine users consistently develop working memory (WM) impairments but the mediating molecular mechanisms are unknown so far. Recent evidence suggests that the serotonin (5-HT) system is altered by chronic cocaine use, while also being involved in WM processing. Thus, we investigated the effects of genetic variations impacting 5-HT activity and of peripheral 5-HT transporter (5-HTT) mRNA expression on WM performance in cocaine users and stimulant-naïve controls.

Two-hundred-twenty participants (126 cocaine users, 94 controls) were assessed with visuospatial, spatial and verbal WM tasks, genotyped for the length polymorphism in the promoter region of the 5-HTT (*5-HTTLPR*), the variable number of tandem repeats in the second intron of the 5-HTT (VNTR In2), two single nucleotide polymorphisms (rs4570625, rs1386497) in the tryptophan hydroxylase-2 (TPH2) gene and quantified for peripheral 5-HTT mRNA expression in whole blood samples.

Several significant gene*environment interactions between 5-HT genotypes and cocaine use on WM emerged: In cocaine users, the long/long (*5-HTTLPR*), 9+10/9+10 (VNTR In2) and C/C (TPH2 rs1386497) genotypes were risk-alleles for WM impairments, while in healthy controls these polymorphisms were associated with improved WM performance. Analogously, high 5-HTT mRNA levels were associated with worse executive WM performance in cocaine users but with increased performance in controls.

These gene*environment interactions suggest that the 5-HT system plays an important role in the development of cognitive deficits in chronic cocaine users. Hence, pharmacological compounds targeting 5-HT neurotransmission might be promising for the treatment of cognitive deficits in cocaine dependence.

Introduction

Cocaine users consistently show impairments in several cognitive domains such as attention, working memory (WM), declarative memory, and executive functions (Beveridge *et al*, 2008; Goldstein *et al*, 2004; Vonmoos *et al*, 2013). Among these, WM seems to be affected first and to show greatest deficits after ongoing use (Jovanovski *et al*, 2005; Vonmoos *et al*, 2013). Thus, WM was claimed to be particularly susceptible to cocaine consumption (Robbins and Arnsten, 2009; Vonmoos *et al*, 2013). Recent longitudinal data furthermore suggest that WM deficits in cocaine users are likely drug-induced rather than predisposed (Vonmoos *et al*, 2014). However, the molecular mechanism by which such cognitive deficits develop, remains unclear so far.

Increasing evidence from rodent and human studies suggests that the serotonin (5-HT) system plays an important role in WM processing (Fitzgerald, 2011). For instance, altering 5-HT synthesis by tryptophan depletion revealed that 5-HT transmission positively correlates with activity in the prefrontal cortex (PFC) and with WM performance (Allen *et al*, 2006; Porter *et al*, 2003). These effects are assumed to be mediated by 5-HT_{2A} and 5-HT_{1A} receptors in the PFC (Puig and Gullledge, 2011; Williams *et al*, 2002). Moreover, in genetic studies, the length polymorphism (short vs. long allele) in the promoter region of the serotonin transporter (5-HTT) gene (*5-HTTLPR*, *SLC6A4*) and more recently a single nucleotide polymorphism (SNP, rs4570625) in the promoter region of the tryptophan hydroxylase 2 (TPH2) gene have been found to influence WM performance (Bosia *et al*, 2010; Homberg and Lesch, 2011; Reuter *et al*, 2008; Zilles *et al*, 2012). This is plausible because the 5-HTT terminates 5-HT from the synaptic cleft by 5-HT reuptake after synaptic signaling (Glatz *et al*, 2003; Lesch *et al*, 1996) and the TPH2 (the rate-limiting enzyme of 5-HT synthesis) appears to facilitate 5-HT neurotransmission by increasing 5-HT availability (Kim *et al*, 2002). However, results of the genetic studies are inconsistent. Some authors found the long allele of *5-HTTLPR* to be associated with superior WM performance (Bosia *et al*, 2010; Zilles *et al*, 2012), whereas others claimed the short allele to be associated with improved performance

(Anderson *et al*, 2012; Enge *et al*, 2011). It was argued that these conflicting results may either be due to a task-specificity of the 5-HTTLPR (Anderson *et al*, 2012; Zilles *et al*, 2012), or due to undetected and moderating environmental factors (Zilles *et al*, 2012).

The 5-HTTLPR has been intensively investigated regarding its potential impact on the risk for cocaine addiction but the results are mixed showing either an association of the long allele with cocaine dependence only in European patients (Patkar *et al*, 2002) or an association of the short allele with cocaine/heroin dependence in Americans of African descent (Enoch *et al*, 2011) or no significant associations of the 5-HTTLPR with cocaine addiction and treatment outcome (Mannelli *et al*, 2005; Patkar *et al*, 2001; Tristan-Noguero *et al*, 2013). Also polymorphisms in the TPH2 gene have been analyzed but they did not show any associations with cocaine dependence (Dahl *et al*, 2006).

Several rodent studies have shown that 5-HT neurotransmission is drastically increased after acute and reduced after chronic cocaine administration (for a review see: Muller and Homberg, 2015). Positron emission tomography (PET) studies in monkeys showed increased 5-HTT availability in frontal areas after chronic cocaine administration, whereas PET studies in humans revealed that chronic use is associated with reduced 5-HT_{1B} receptor binding in the PFC (Gould *et al*, 2011; Matuskey *et al*, 2013). In addition, studies in rats demonstrated that the TPH2 activity in the raphe nucleus increases after chronic cocaine administration (Vrana *et al*, 1993). In their comprehensive review on the role of 5-HT in drug addiction, Müller and Homberg (2015) concluded that chronic cocaine use might induce an increase in terminal 5-HTT function with reduced extracellular 5-HT levels in cortical areas. Subsequent acute cocaine challenges may then encounter more terminal and less somatodendritic 5-HTT targets and, thus, induce an intensified cortical 5-HT increase further reiterating the change in 5-HT function.

The aim of this study was to investigate whether 5-HTT and TPH2 polymorphisms are associated with WM deficits in cocaine users. As outlined above, polymorphisms in 5-HTT and TPH2 genes (Bosia *et al*, 2010; Homberg and Lesch, 2011; Reuter *et al*, 2008; Zilles *et al*, 2012) as well as chronic cocaine use

(Muller and Homberg, 2015) are assumed to impact 5-HT neurotransmission. Given that the 5-HT system has been linked to WM performance (Allen *et al*, 2006; Porter *et al*, 2003), we hypothesized that 5-HTT and TPH2 polymorphisms interact with cocaine use regarding WM performance. In addition to the above-mentioned 5-HTTLPR and the TPH2 polymorphism (rs4570625), we further examined the variable number tandem repeats polymorphism within intron 2 of the 5-HTT (VNTR In2) and a second promising SNP in the TPH2 gene (rs1386497). The VNTR In2 was chosen because it is assumed to have 5-HTT transcription regulating properties and the rs1386497 was selected due to its association with cognitive deficits in attention deficit hyperactivity disorder (ADHD) (MacKenzie and Quinn, 1999; Manor *et al*, 2008). Besides these polymorphisms, we measured 5-HTT (SLC6A4) mRNA expression in peripheral blood in order to detect potential interactions between 5-HTT and TPH2 genotype, 5-HTT gene expression and cocaine use. Finally, because of the inconsistent results of previous studies, we used various WM tasks in order to compare genotype effects for visuospatial, spatial, and verbal WM subdomains.

Materials and Methods

Participants

A total sample of 220 participants (126 chronic cocaine users and 94 stimulant-naïve controls) was investigated as part of the longitudinal Zurich Cocaine Cognition Study (ZuCo²St: (Vonmoos *et al*, 2013; Vonmoos *et al*, 2014)). Cocaine users were included if they indicated cocaine as their primary drug of choice, showed a use of >0.5g/month, and a maximum abstinence duration no longer than 6 months. Exclusion criteria for all participants were any current or previous neurological disorders or head injuries and use of prescription drugs affecting cognitive functioning. Additional exclusion criteria for the cocaine users were regular use of opioids or a polytoxic drug use pattern according to DSM-IV as well as any current or previous Axis-I DSM-IV psychiatric disorders with exception of cocaine and alcohol abuse/dependence, a history of depression, and ADHD. Specific exclusion criteria for the control group were regular drug use (lifetime >15 occasions) with the exception of nicotine dependence and occasional cannabis use and any current or previous Axis-I DSM-IV psychiatric disorder (American Psychiatric Association, 1994). Participants self-reports regarding drug use were controlled by toxicological analyses of urine and 6cm (if possible) hair samples (see also Table S1). More detailed information about selection, recruitment, and drug screenings have been presented in detail in our previous work (Hulka *et al*, 2014; Preller *et al*, 2013; Vonmoos *et al*, 2013). The presented study was approved by the Cantonal Ethics Committee of Zurich. All participants provided written informed consent prior to the study and were financially compensated for their participation.

Procedure

All participants were examined by trained psychologists using a Structural Clinical Interview (SCID-I)(Wittchen *et al*, 1997) according to DSM-IV. Drug use was assessed by using a structured and

standardized Interview for Psychotropic Drug Consumption (Quednow *et al*, 2004). Subsequently, participants underwent a battery of neuropsychological evaluations including the Spatial Working Memory task (SWM) and the Paired Associates Learning task (PAL) from the Cambridge Neuropsychological Test Automated Battery (CANTAB, www.cambridgecognition.com)(Strauss *et al*, 2006) as well as the Letter Number Sequencing Task (LNST) (Wechsler, 1997). In order to draw conclusions about global WM processes, the three main parameters of these tests (SWM total errors, PAL first trial memory score, LNST correct responses) were z-transformed (based on means and standard deviations of the control group) and incorporated into a global WM capacity index (as described in our previous work: Vonmoos *et al*, 2013). To additionally investigate an executive WM component, we separately analyzed the SMW strategy score that is thought to primarily reflect executive functioning, i.e. strategic planning (Owen *et al*, 1990).

Genotyping and mRNA Quantification

The *5-HTTLPR*, VNTR In2, and TPH2 polymorphisms were determined as described in the Supplemental Material. The *5-HTTLPR* was assessed in its bi-allelic (short vs. long allele) instead of its tri-allelic form (LA vs. LG, SA and SG, see: Hu *et al*, 2006) in order to compare our results with previous studies on the effects of *5-HTTLPR* on WM. Furthermore, subjects with 9 copies (2.7%) in the VNTR In2 were pooled with subjects carrying 10 copies resulting in three genotype groups: 9+10/9+10 vs. 9+10/12 vs. 12/12. Also, subjects with the rare T/T genotype (6,2%) of TPH2 rs4570625 and the rare C/C genotype (3.6%) in TPH2 rs1386497 were pooled with subject with the G/T and the A/C genotype, respectively, resulting in the two genotype groups: G/G vs. G+T/T and A/A vs. A+C/C.

Total RNA was isolated from whole blood using the RNA isolation NucleoSpin RNA Blood in combination with the NucleoSpin RNA/DNA Buffer Set according to the manufacturer's recommendation (Macherey-Nagel AG, Oersingen, Switzerland). Total RNA samples were spectrophotometrically scanned (NanoVue,

GE Healthcare Life Sciences) to obtain the A260/A280 of >1.9 and concentration levels. Additionally, for RNA quality resulting with RIN (RNA integrity number) values, all samples were measured on the automated electrophoresis system (Experion, BioRad Co., Hercules, CA, USA). Quantitative real-time RT-PCR was conducted for 5-HTT (*SLC6A4*) and six additional reference genes (*ACTB*, *GAPDH*, *ALAS1*, *RPL13A*, *PPIA*, and 18S ribosomal RNA) as described previously (Grunblatt *et al*, 2009). Total RNA (500 ng) from each sample was reverse transcribed using iScript cDNA synthesis kit (BioRad Co., Hercules, CA, USA). Each amplification was performed in a total volume of 20µl containing 5µl QuantiTect SYBR Green PCR kit (Qiagen) and the specific primer mix (PrimerAssay- Qiagen). PCR conditions were run according to manufacturer's manual (Qiagen). A melting point analysis was conducted for each assay to confirm specificity of PCR products and all PCR reactions were run in triplicates. The program LinRegPCR (www.hartfaalcentrum.nl) was used to determine the PCR efficiency. Gene expression and normalization analysis with the most stable reference genes was conducted using the QBase plus software (Biogazelle) (Vandesompele *et al*, 2002). The software detected that the reference gene *RPL13A* was least stable and therefore this gene was excluded and normalization analysis was conducted using the five other reference genes. We also tried to determine TPH2 mRNA but the melting point analysis revealed an insufficient quality of the PCR products indicating that the TPH2 mRNA was degraded.

Statistical Analyses

Global and task-specific WM effects were analyzed separately for each polymorphism using analyses of variance (ANOVA) with the factors: *group* (cocaine users vs. controls) and *genotype* (5-HTTLPR and VNTR In2: threefold, TPH2 rs4570625 and rs1386497: twofold, see above). Additionally, analyses of covariance (ANCOVA) with the covariates *age*, *years of education*, *BDI*, and *ADHD* were calculated to investigate possible influences of these factors. The influence of 5-HTT mRNA expression on WM performance was investigated using Pearson correlations, whereas mRNA expression*genotype interactions were

examined using ANCOVAs with *mRNA expression* as the covariate. Results were considered significant if $p < .05$ after correction for multiple comparisons using the Hochberg method (Hochberg, 1988).

Hochberg's procedure (a step-up modification of the Bonferroni method) tests each partition hypothesis using all the order statistics by formulating a sequence of critical values based on Simes' inequality.

Statistical analyses were conducted with SPSS (Version 20.0), testing of linkage disequilibrium (LD)

between polymorphisms was performed with the software *Haploview* (Barrett *et al*, 2005) and

associations between polymorphisms and cocaine use was analyzed with the Armitage's Trend Test.

Results

Demography and association analysis

Cocaine users and stimulant-naïve controls did not differ regarding age and sex distribution. However, cocaine users had significantly fewer years of education, higher scores in the Beck Depression Inventory (BDI) and higher scores of ADHD than controls (see Table 1). Beyond these group differences, we did not find any significant *genotype* or *group*genotype* interaction effects on BDI (all $p>.05$) or ADHD (all $p>.05$). For additional information on the substance use of both groups see Table S1 in the Supplemental Material. Genotype frequencies of all four polymorphisms (see also Table S2) were distributed in accordance with the Hardy-Weinberg Equilibrium (HWE) and the SNPs rs4570625 and rs1386497 were found to be in minimal Linkage Disequilibrium (LD, $D'=1.0$, LOD: 5.16, $r^2=0.06$). None of the polymorphisms was associated with cocaine use per se (Table S2) and within the group of cocaine users genotype groups did not differ regarding cocaine consumption (Table S3).

Global WM Capacity

Global WM performance was affected by a significant main effect of the factor *group*, whereas the factor *genotype* alone was not significant in any analysis (Table 2). However, there were significant *group*genotype* interactions for all variants except for TPH2 rs4570625, which nonetheless showed a similar trend. Cocaine users revealed a worse WM performance compared to controls as shown previously for this sample (Vonmoos *et al*, 2013), whereby 5-HTTLPR, VNTR In2, and TPH2 rs1386497 *genotype* inversely affected WM in cocaine users and controls (Figure 1). These effects remained significant after correction for multiple comparisons (see Table 2) and even when *age*, *years of education*, *BDI* and *ADHD* were introduced as covariates (5-HTTLPR: $F(2,207)=5.28$, $p_{cor}<.05$; VNTR In2: $F(2,202)=7.29$, $p_{cor}<.01$; TPH2 rs1386497: $F(1,207)=5.14$, $p_{cor}<.05$). Among the covariates, *age* showed a

significant main effect in the ANCOVAs of all three significant polymorphisms (all $p_{cor} < .001$), whereas *years of education* was only significant for 5-HTTLPR and TPH2 rs1386497 (both $p_{cor} < .05$).

WM Subdomains and Executive WM

Table 2 further reveals that the above described main effects of *group* and the *group*genotype* interactions were also found within the spatial, verbal, and visuospatial WM subdomains and in the executive WM parameter *SMW strategy*. In *SWM strategy*, we observed similar significant *group*genotype* interactions with 5-HTTLPR and VNTR In2 genotype as shown for global WM above (see Figure S1). Regarding the spatial WM subdomain (CANTAB SWM total errors), significant interactions (corrected for multiple testing) were observed for VNTR In2 and TPH2 rs1386497 genotypes while 5-HTTLPR and TPH2 rs4570625 displayed strong trends. LNST also showed moderate interaction effects of 5-HTTLPR, VNTR In2 genotypes and TPH2 rs1386497 with group, which remained trends after multiple-test correction. Finally, in CANTAB PAL, no interaction effects were strong enough to survive correction for multiple testing. It should be noted that all subdomains (with exception of PAL regarding TPH2 rs1386497) showed the same pattern of *group*genotype* interaction as illustrated for the global WM capacity index in Figure 1 (see Figure S2).

5-HTT mRNA expression

Peripheral mRNA expression did not differ between controls and cocaine users and across genotype groups (see Table S4). Also, correlations between 5-HTT mRNA expression and WM performance did not reveal any significant results, neither in the total sample nor in both subgroups. However, using a median split of the mRNA expression (high vs. low expression) together with the factor *group* in a two-way ANOVA revealed a significant *group*mRNA expression* interaction ($F(1,107)=6.36$, $p_{cor} > .05$, corrected for *age* and *years of education*) in SWM strategy (Figure 2). Additionally, by including *mRNA expression* as a

covariate in an ANCOVA together with *group* and *genotype*, we found significant three-way interactions of *group*genotype*mRNA expression* for VNTR In2 in the global WM capacity index ($F(4, 96)=5.04$, $p_{cor}<.01$), in LNST ($F(4, 96)=3.93$, $p_{cor}<.05$) and in PAL ($F(4, 96)=3.70$, $p_{cor}<.05$), for TPH2 rs4570625 in the global WM capacity index ($F(2, 101)=9.15$, $p_{cor}<.001$), in LNST ($F(2, 101)=4.85$, $p_{cor}<.05$), in PAL ($F(2, 101)=7.16$, $p_{cor}<.01$) and in SWM strategy ($F(2, 101)=5.51$, $p_{cor}<.05$) and for TPH2 rs1386497 in LNST ($F(1, 207)=4.401$, $p_{cor}<.05$). These threefold interactions (see Figure S3-S7) were mainly explained by a general interaction between *gene expression* and *group* mirroring the *group*genotype* interaction shown above (see Figure 2).

Discussion

The present study was designed to investigate whether WM impairments in cocaine users are mediated by polymorphisms within the 5-HT system. We demonstrated that three of four investigated 5-HT-impacting polymorphisms (*5-HTTLPR*, VNTR In2, TPH2 rs1386497) mediate WM performance and that their effects on WM depend on cocaine use of participants.

Concerning the *5-HTTLPR* polymorphism, we found the long/long genotype to be a risk factor for the presence of WM impairments in cocaine users, while it was associated with improved WM performance in healthy controls. Even though, the effects of *5-HTTLPR* on 5-HT neurotransmission in humans is a subject of controversial debate, our results can be interpreted in consideration of previous animal studies. Carriers of the short allele of the *5-HTTLPR* polymorphism are assumed to display decreased cerebral 5-HTT expression and thus decreased 5-HT reuptake after synaptic signaling (Glatz *et al*, 2003; Lesch *et al*, 1996). However, carrying the short allele does not appear to result in increased 5-HT neurotransmission. In fact, increasing evidence from studies using 5-HTT knock-out mice suggests that having the short allele results in decreased 5-HT neurotransmission based on long-term developmental effects of reduced 5-HT reuptake on cell number, firing rate, and postsynaptic receptor sensitivity (Kim *et al*, 2005; Lira *et al*, 2003). Assuming, thus, that healthy control subjects carrying the short/short genotype have lower 5-HT activity and a reduced clearance rate after activation than long/long controls, our results are consistent with previous studies showing decreased cognitive functioning in participants with lower 5-HT levels (Allen *et al*, 2006; Fitzgerald, 2011; Murphy *et al*, 2002; Porter *et al*, 2003; Rogers *et al*, 2003). However, this interpretation is speculative because there is no consensus on the effects of *5-HTTLPR* on 5-HT neurotransmission in humans so far. For instance, two molecular imaging studies employing PET in humans did not find an association between the *5-HTTLPR* genotype and 5-HTT binding (Murthy *et al*, 2010; Shioe *et al*, 2003), while two other PET studies showed increased 5-HTT densities in

putamen and midbrain, respectively, of long allele carriers (Praschak-Rieder *et al*, 2007; Reimold *et al*, 2007).

In cocaine users, carriers of the short/short genotype showed comparable WM performance as short/short genotype carriers of the controls. This finding is plausible considering the effects of cocaine use on the 5-HT system. Chronic cocaine administration is postulated (Muller and Homberg, 2015) to simultaneously result in an overexpression of 5-HTT at terminal sites of neurons and in an attenuated expression of 5-HTT at somatodendritic sites (Gould *et al*, 2011; Muller *et al*, 2002; Muller and Homberg, 2015). By this, chronic cocaine consumption is thought to result in a chronically reduced 5-HT neuronal activity together with a more efficient removal of 5-HT from the synaptic cleft in terminal areas of the 5-HT projections. The *5-HTTLPR* long/long genotype probably displays more 5-HTT under basal conditions, which may offer more target sites for cocaine interactions. One may, therefore, speculate that 5-HTT adaptations after chronic cocaine use are stronger in the long/long than in the short/short genotype. This may result in a stronger reduction of basal 5-HT activity in the long/long genotype and, thus, in more pronounced WM deficits. It could thus be assumed that, under such conditions, having a genetically limited 5-HTT expression protects against cocaine-induced adaptations of the 5-HTT function. Hence, this view is supported by our data indicating unaffected WM performance in cocaine users carrying the short allele.

Our results regarding the *5-HTTLPR* polymorphism seem to confirm the hypothesis that increased 5-HT activity is associated with improved cognitive functioning (in our case WM) in healthy controls (Murphy *et al*, 2002; Rogers *et al*, 2003). Moreover, we went one step further by showing that environmental factors (such as cocaine use) are able to moderate the direction of the polymorphism effects. This finding is especially interesting considering the conflicting results of previous studies suggesting that the effects of the *5-HTTLPR* might be task-specific (Anderson *et al*, 2012; Zilles *et al*, 2012). However, we were able to demonstrate consistent genotype effects across WM tasks measuring spatial (SWM), verbal (LNST),

and visuospatial (PAL) WM subdomains (see Table 1 and Figures S2). Hence, we propose that undetected gene*environment interactions (such as the discovered genotype-cocaine interaction) could provide an explanation for the previously conflicting results.

Our findings are further supported by the 5-HTT mRNA expression results. Analyzing peripheral mRNA expression in whole blood samples, we found a similar *group*genotype* interaction effect of 5-HTTLPR in form of an interaction of *group*mRNA expression* also indicating higher WM performance in controls with increased 5-HTT expression and lower WM performance in cocaine users with increased 5-HTT expression. This is consistent with previous studies showing that 5-HTT mRNA expression can be used to replicate 5-HTTLPR genotype findings (Gardner *et al*, 2009).

Regarding the two SNPs in the TPH2 gene, we were not able to clearly replicate previous studies postulating the rare T/T genotype of rs4570625 to be associated with lower WM performance (Reuter *et al*, 2008). However, we found a significant interaction between TPH2 rs1386497 (which is in LD with rs4570625) and cocaine use. The A/A genotype of rs1386497 was associated with lower performance in cocaine users, whereas healthy controls carrying the A/A genotype showed higher WM performance. Tryptophan hydroxylase is a rate-limiting enzyme of the 5-HT synthesis (Kim *et al*, 2002) and although the rs1386497 has not yet been reported in relation to WM, its A allele has been associated with ADHD and cognitive deficits in ADHD (Manor *et al*, 2008; Sheehan *et al*, 2005). Interestingly, we reported in our previous work that the co-occurrence of cocaine consumption and ADHD had a particularly negative impact on cognitive functioning (Vonmoos *et al*, 2013). However, including ADHD as covariate still revealed an ADHD-independent interaction effect of TPH2 rs1386497 and cocaine use on WM performance.

Concerning the VNTR In2, we demonstrated that controls revealed higher WM performance when carrying the short allele (9+10/9+10 copies) compared with carriers of the long allele (12/12 copies). In cocaine users, again, this pattern was inverted. It has been proposed that the VNTR In2 polymorphism

acts as a transcriptional regulator of the 5-HTT expression but its exact effect is not fully clear yet (MacKenzie and Quinn, 1999). Comparing our findings of the 5-HTTLPR with the findings of the VNTR In2 reveals that the long/long genotype of *5-HTTLPR* seems to have a similar effect on WM as the 12 copies variant of the VNTR In2. This specific pattern has already been found by other genetic studies with psychiatric patients suffering from personality and psychosomatic disorders (Garcia *et al*, 2010; Kirtak *et al*, 2008).

In sum, we presented evidence that genetic variations, putatively influencing 5-HT activity, mediate the development of WM impairments in cocaine users and that increased 5-HTT expression in the periphery goes along with these WM deficits. Our results support the view that chronic cocaine use enhances terminal 5-HTT function and that this enhanced activity, together with adaptations of postsynaptic 5-HT receptors, is responsible for the cognitive decline seen in cocaine users (Muller and Homberg, 2015). Alternatively, it might be speculated that specific 5-HTT polymorphisms, such as *5-HTTLPR* and VNTR In2, impact the protein tertiary structure of the 5-HTT leading to a changed affinity to cocaine. In that case, genetic variations could also impact the acute effects of cocaine on 5-HT reuptake inhibition. Future studies should therefore investigate whether there are genotype differences in the acute effects of cocaine administration on a neuropharmacological as well as on a behavioral level. In addition, functional and molecular imaging techniques might be a promising approach to get a deeper understanding of the mechanisms underlying the cocaine-genotype interaction on WM.

In conclusion, we demonstrated that three 5-HT-impacting polymorphisms are risk-alleles for the development of WM deficits in cocaine users. Hence, pharmacological compounds targeting 5-HT neurotransmission might be a promising treatment option of WM deficits in cocaine addiction. Additionally, our results regarding the *5-HTTLPR* and the VNTR In2 polymorphisms in healthy controls seem to confirm the hypothesis (Fitzgerald, 2011) that increased 5-HT activity improves cognitive functioning (in our case WM). Finally, the discovered gene*environment interaction (i.e. cocaine use

moderated genotype effects) indicates that the complexity of the 5-HT systems requires us to discover other relevant factors to fully understand the involvement of 5-HT in WM processing.

Funding and Disclosure

The study was supported by grants from the Swiss National Science Foundation (SNSF; grant No. PP00P1-123516/1 and PP00P1-146326/1) and the Olga Mayenfisch Foundation. None of the authors declared any conflict of interest.

Acknowledgments

We would like to thank Vera Guttenthaler and Miryame Hofmann for technical assistance.

References

- Allen PP, Cleare AJ, Lee F, Fusar-Poli P, Tunstall N, Fu CH, *et al* (2006). Effect of acute tryptophan depletion on pre-frontal engagement. *Psychopharmacology (Berl)* **187**(4): 486-497.
- American Psychiatric Association (1994). *Diagnostic and Statistical Manual of Mental Disorders (4th edn) (DSM-IV)* APA.
- Anderson DE, Bell TA, Awh E (2012). Polymorphisms in the 5-HTTLPR gene mediate storage capacity of visual working memory. *J Cogn Neurosci* **24**(5): 1069-1076.
- Barrett JC, Fry B, Maller J, Daly MJ (2005). Haploview: analysis and visualization of LD and haplotype maps. *Bioinformatics* **21**(2): 263-265.
- Beveridge TJ, Gill KE, Hanlon CA, Porrino LJ (2008). Review. Parallel studies of cocaine-related neural and cognitive impairment in humans and monkeys. *Philos Trans R Soc Lond B Biol Sci* **363**(1507): 3257-3266.
- Bosia M, Anselmetti S, Pirovano A, Ermoli E, Marino E, Bramanti P, *et al* (2010). HTTLPR functional polymorphism in schizophrenia: executive functions vs. sustained attention dissociation. *Prog Neuropsychopharmacol Biol Psychiatry* **34**(1): 81-85.
- Dahl JP, Cubells JF, Ray R, Weller AE, Lohoff FW, Ferraro TN, *et al* (2006). Analysis of variations in the tryptophan hydroxylase-2 (TPH2) gene in cocaine dependence. *Addict Biol* **11**(1): 76-83.
- Enge S, Fleischhauer M, Lesch KP, Reif A, Strobel A (2011). Serotonergic modulation in executive functioning: linking genetic variations to working memory performance. *Neuropsychologia* **49**(13): 3776-3785.
- Enoch MA, Gorodetsky E, Hodgkinson C, Roy A, Goldman D (2011). Functional genetic variants that increase synaptic serotonin and 5-HT₃ receptor sensitivity predict alcohol and drug dependence. *Mol Psychiatry* **16**(11): 1139-1146.
- Fitzgerald PJ (2011). A neurochemical yin and yang: does serotonin activate and norepinephrine deactivate the prefrontal cortex? *Psychopharmacology (Berl)* **213**(2-3): 171-182.
- Garcia LF, Aluja A, Fibla J, Cuevas L, Garcia O (2010). Incremental effect for antisocial personality disorder genetic risk combining 5-HTTLPR and 5-HTTVNTR polymorphisms. *Psychiatry Res* **177**(1-2): 161-166.
- Gardner KL, Hale MW, Lightman SL, Plotsky PM, Lowry CA (2009). Adverse early life experience and social stress during adulthood interact to increase serotonin transporter mRNA expression. *Brain Res* **1305**: 47-63.

Glatz K, Mossner R, Heils A, Lesch KP (2003). Glucocorticoid-regulated human serotonin transporter (5-HTT) expression is modulated by the 5-HTT gene-promotor-linked polymorphic region. *J Neurochem* **86**(5): 1072-1078.

Goldstein RZ, Leskovjan AC, Hoff AL, Hitzemann R, Bashan F, Khalsa SS, *et al* (2004). Severity of neuropsychological impairment in cocaine and alcohol addiction: association with metabolism in the prefrontal cortex. *Neuropsychologia* **42**(11): 1447-1458.

Gould RW, Gage HD, Banks ML, Blaylock BL, Czoty PW, Nader MA (2011). Differential effects of cocaine and MDMA self-administration on cortical serotonin transporter availability in monkeys. *Neuropharmacology* **61**(1-2): 245-251.

Grunblatt E, Bartl J, Zehetmayer S, Ringel TM, Bauer P, Riederer P, *et al* (2009). Gene expression as peripheral biomarkers for sporadic Alzheimer's disease. *J Alzheimers Dis* **16**(3): 627-634.

Hochberg Y (1988). A sharper Bonferroni procedure for multiple tests of significance. *Biometrika* **75**(4): 3.

Homberg JR, Lesch KP (2011). Looking on the bright side of serotonin transporter gene variation. *Biol Psychiatry* **69**(6): 513-519.

Hu XZ, Lipsky RH, Zhu G, Akhtar LA, Taubman J, Greenberg BD, *et al* (2006). Serotonin transporter promoter gain-of-function genotypes are linked to obsessive-compulsive disorder. *Am J Hum Genet* **78**(5): 815-826.

Hulka LM, Eisenegger C, Preller KH, Vonmoos M, Jenni D, Bendrick K, *et al* (2014). Altered social and non-social decision-making in recreational and dependent cocaine users. *Psychol Med* **44**(5): 1015-1028.

Jovanovski D, Erb S, Zakzanis KK (2005). Neurocognitive deficits in cocaine users: a quantitative review of the evidence. *J Clin Exp Neuropsychol* **27**(2): 189-204.

Kim DK, Tolliver TJ, Huang SJ, Martin BJ, Andrews AM, Wichems C, *et al* (2005). Altered serotonin synthesis, turnover and dynamic regulation in multiple brain regions of mice lacking the serotonin transporter. *Neuropharmacology* **49**(6): 798-810.

Kim SW, Park SY, Hwang O (2002). Up-regulation of tryptophan hydroxylase expression and serotonin synthesis by sertraline. *Mol Pharmacol* **61**(4): 778-785.

Kirtak N, Inaloz HS, Akcali C, Erdal E, Herken H, Yildirim M, *et al* (2008). Association of serotonin transporter gene-linked polymorphic region and variable number of tandem repeat polymorphism of the serotonin transporter gene in lichen simplex chronicus patients with psychiatric status. *Int J Dermatol* **47**(10): 1069-1072.

Lesch KP, Bengel D, Heils A, Sabol SZ, Greenberg BD, Petri S, *et al* (1996). Association of anxiety-related traits with a polymorphism in the serotonin transporter gene regulatory region. *Science* **274**(5292): 1527-1531.

Lira A, Zhou M, Castanon N, Ansorge MS, Gordon JA, Francis JH, *et al* (2003). Altered depression-related behaviors and functional changes in the dorsal raphe nucleus of serotonin transporter-deficient mice. *Biol Psychiatry* **54**(10): 960-971.

MacKenzie A, Quinn J (1999). A serotonin transporter gene intron 2 polymorphic region, correlated with affective disorders, has allele-dependent differential enhancer-like properties in the mouse embryo. *Proc Natl Acad Sci U S A* **96**(26): 15251-15255.

Mannelli P, Patkar AA, Murray HW, Certa K, Peindl K, Mattila-Evenden M, *et al* (2005). Polymorphism in the serotonin transporter gene and response to treatment in African American cocaine and alcohol-abusing individuals. *Addict Biol* **10**(3): 261-268.

Manor I, Laiba E, Eisenberg J, Meidad S, Lerer E, Israel S, *et al* (2008). Association between tryptophan hydroxylase 2, performance on a continuance performance test and response to methylphenidate in ADHD participants. *Am J Med Genet B Neuropsychiatr Genet* **147B**(8): 1501-1508.

Matuskey D, Bhagwagar Z, Planeta B, Pittman B, Gallezot JD, Chen J, *et al* (2013). Reductions in Brain 5-HT Receptor Availability in Primarily Cocaine-Dependent Humans. *Biol Psychiatry* **76**(10): 816-822.

Muller CP, De Souza Silva MA, DePalma G, Tomaz C, Carey RJ, Huston JP (2002). The selective serotonin(1A)-receptor antagonist WAY 100635 blocks behavioral stimulating effects of cocaine but not ventral striatal dopamine increase. *Behav Brain Res* **134**(1-2): 337-346.

Muller CP, Homberg JR (2015). The role of serotonin in drug use and addiction. *Behav Brain Res* **277**: 146-192.

Murphy FC, Smith KA, Cowen PJ, Robbins TW, Sahakian BJ (2002). The effects of tryptophan depletion on cognitive and affective processing in healthy volunteers. *Psychopharmacology (Berl)* **163**(1): 42-53.

Murthy NV, Selvaraj S, Cowen PJ, Bhagwagar Z, Riedel WJ, Peers P, *et al* (2010). Serotonin transporter polymorphisms (SLC6A4 insertion/deletion and rs25531) do not affect the availability of 5-HTT to [11C] DASB binding in the living human brain. *Neuroimage* **52**(1): 50-54.

Owen AM, Downes JJ, Sahakian BJ, Polkey CE, Robbins TW (1990). Planning and spatial working memory following frontal lobe lesions in man. *Neuropsychologia* **28**(10): 1021-1034.

Patkar AA, Berrettini WH, Hoehe M, Hill KP, Gottheil E, Thornton CC, *et al* (2002). No association between polymorphisms in the serotonin transporter gene and susceptibility to cocaine dependence among African-American individuals. *Psychiatr Genet* **12**(3): 161-164.

Patkar AA, Berrettini WH, Hoehe M, Hill KP, Sterling RC, Gottheil E, *et al* (2001). Serotonin transporter (5-HTT) gene polymorphisms and susceptibility to cocaine dependence among African-American individuals. *Addict Biol* **6**(4): 337-345.

Porter RJ, Lunn BS, O'Brien JT (2003). Effects of acute tryptophan depletion on cognitive function in Alzheimer's disease and in the healthy elderly. *Psychol Med* **33**(1): 41-49.

Praschak-Rieder N, Kennedy J, Wilson AA, Hussey D, Boovariwala A, Willeit M, *et al* (2007). Novel 5-HTTLPR allele associates with higher serotonin transporter binding in putamen: a [(11)C] DASB positron emission tomography study. *Biol Psychiatry* **62**(4): 327-331.

Preller KH, Ingold N, Hulka LM, Vonmoos M, Jenni D, Baumgartner MR, *et al* (2013). Increased sensorimotor gating in recreational and dependent cocaine users is modulated by craving and attention-deficit/hyperactivity disorder symptoms. *Biol Psychiatry* **73**(3): 225-234.

Puig MV, Gullledge AT (2011). Serotonin and prefrontal cortex function: neurons, networks, and circuits. *Mol Neurobiol* **44**(3): 449-464.

Quednow BB, Kuhn KU, Hoenig K, Maier W, Wagner M (2004). Prepulse inhibition and habituation of acoustic startle response in male MDMA ('ecstasy') users, cannabis users, and healthy controls. *Neuropsychopharmacology* **29**(5): 982-990.

Reimold M, Smolka MN, Schumann G, Zimmer A, Wrase J, Mann K, *et al* (2007). Midbrain serotonin transporter binding potential measured with [(11)C]DASB is affected by serotonin transporter genotype. *J Neural Transm* **114**(5): 635-639.

Reuter M, Esslinger C, Montag C, Lis S, Gallhofer B, Kirsch P (2008). A functional variant of the tryptophan hydroxylase 2 gene impacts working memory: a genetic imaging study. *Biol Psychol* **79**(1): 111-117.

Robbins TW, Arnsten AF (2009). The neuropsychopharmacology of fronto-executive function: monoaminergic modulation. *Annu Rev Neurosci* **32**: 267-287.

Rogers RD, Tunbridge EM, Bhagwagar Z, Drevets WC, Sahakian BJ, Carter CS (2003). Tryptophan depletion alters the decision-making of healthy volunteers through altered processing of reward cues. *Neuropsychopharmacology* **28**(1): 153-162.

Sheehan K, Lowe N, Kirley A, Mullins C, Fitzgerald M, Gill M, *et al* (2005). Tryptophan hydroxylase 2 (TPH2) gene variants associated with ADHD. *Mol Psychiatry* **10**(10): 944-949.

Shioe K, Ichimiya T, Suhara T, Takano A, Sudo Y, Yasuno F, *et al* (2003). No association between genotype of the promoter region of serotonin transporter gene and serotonin transporter binding in human brain measured by PET. *Synapse* **48**(4): 184-188.

Strauss E, Sherman EM, Spreen O (2006). *A Compendium of Neuropsychological Tests: Administration, Norms, and Commentary*, Third Edition edn. Oxford University Press.

Tristan-Noguero A, Fernandez-Castillo N, Roncero C, Sanchez-Mora C, Ramos-Quiroga JA, Daigre C, *et al* (2013). Lack of association between the LPR and VNTR polymorphisms of the serotonin transporter gene and cocaine dependence in a Spanish sample. *Psychiatry Res* **210**(3): 1287-1289.

Vandesompele J, De Preter K, Pattyn F, Poppe B, Van Roy N, De Paepe A, *et al* (2002). Accurate normalization of real-time quantitative RT-PCR data by geometric averaging of multiple internal control genes. *Genome Biol* **3**(7): RESEARCH0034.

Vonmoos M, Hulka LM, Preller KH, Jenni D, Baumgartner MR, Stohler R, *et al* (2013). Cognitive dysfunctions in recreational and dependent cocaine users: role of attention-deficit hyperactivity disorder, craving and early age at onset. *Br J Psychiatry* **203**(1): 35-43.

Vonmoos M, Hulka LM, Preller KH, Minder F, Baumgartner MR, Quednow BB (2014). Cognitive impairment in cocaine users is drug-induced but partially reversible: evidence from a longitudinal study. *Neuropsychopharmacology* **39**(9): 2200-2210.

Vrana SL, Vrana KE, Koves TR, Smith JE, Dworkin SI (1993). Chronic cocaine administration increases CNS tyrosine hydroxylase enzyme activity and mRNA levels and tryptophan hydroxylase enzyme activity levels. *J Neurochem* **61**(6): 2262-2268.

Wechsler DA (1997). *Wechsler Memory Scale (3rd edn) Manual* Psychological Corporation.

Williams GV, Rao SG, Goldman-Rakic PS (2002). The physiological role of 5-HT_{2A} receptors in working memory. *J Neurosci* **22**(7): 2843-2854.

Wittchen HU, Wunderlich U, Gruschwitz S, Zaudig M (1997). *SKID-I. Strukturiertes Klinisches Interview für DSM-IV Achse I: Psychische Störungen. [SCID-I. Structured Clinical Interview for DSM-IV Axis I Disorders]* Hogrefe.

Zilles D, Meyer J, Schneider-Axmann T, Ekawardhani S, Gruber E, Falkai P, *et al* (2012). Genetic polymorphisms of 5-HTT and DAT but not COMT differentially affect verbal and visuospatial working memory functioning. *Eur Arch Psychiatry Clin Neurosci* **262**(8): 667-676.

Tables

Table 1 Demographic data of healthy controls and cocaine users (means and *SD*).

Demographic Data	Controls (<i>n</i> =94)	Cocaine users (<i>n</i> =126)	<i>t</i> -test/ χ^2	<i>df</i>	<i>p</i>
Age, mean (<i>SD</i>)	30.0 (8.8)	30.1 (8.9)	-0.1	218	0.917
Sex (male / female)	68 / 26	91 / 35	0.0	1	0.554
Years of education, mean (<i>SD</i>)	10.7 (1.8)	10.2 (1.7)	2.1	218	0.041
BDI Score, mean (<i>SD</i>)	4.2 (4.0)	8.6 (7.1)	-5.8	204.2	0.000
ADHD DSM-IV Score, mean (<i>SD</i>)	7.9 (5.0)	14.4 (9.4)	-6.5	198.7	0.000

BDI: Beck Depression Inventory, ADHD: Attention Deficit Hyperactivity Disorder

Table 2 Analyses of variance for global working memory capacity, working memory subdomains, and executive working memory across all investigated polymorphisms.

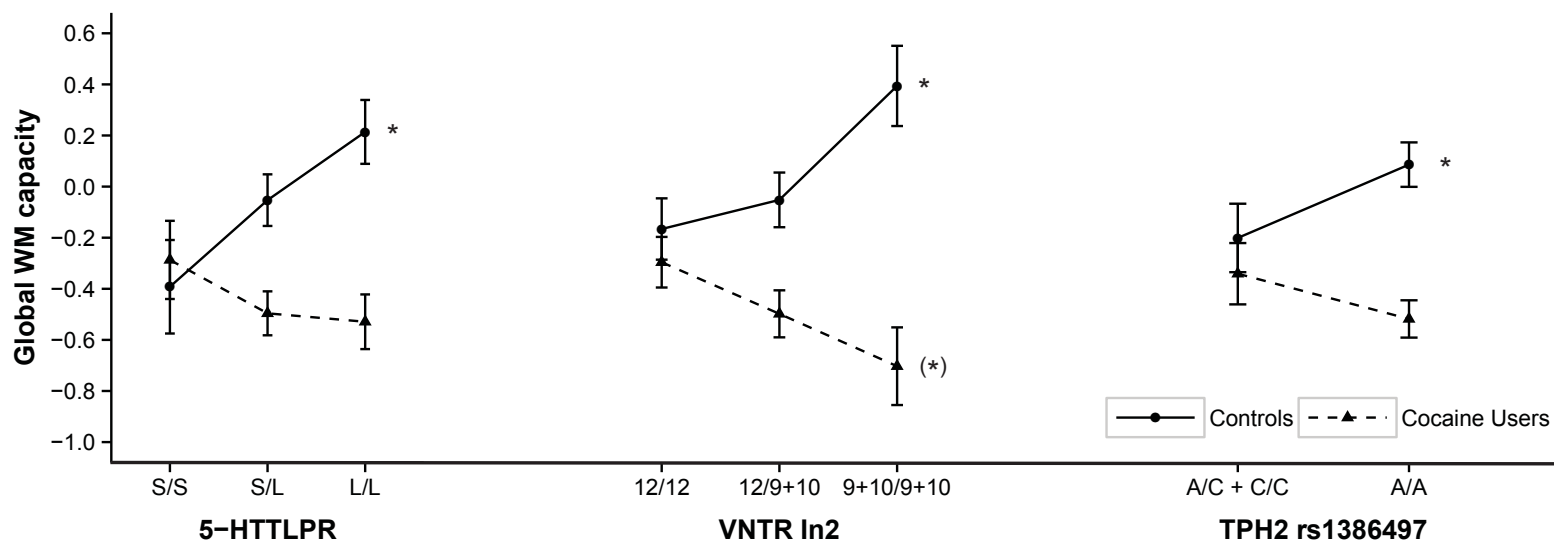
	5-HTTLPR					VNTR In2					TPH2 (rs4570625)					TPH2 (rs1386497)				
	<i>F</i>	<i>df</i>	<i>p_{ncor}</i>	ηp^2	<i>p_{cor}</i>	<i>F</i>	<i>df</i>	<i>p_{ncor}</i>	ηp^2	<i>p_{cor}</i>	<i>F</i>	<i>df</i>	<i>p_{ncor}</i>	ηp^2	<i>p_{cor}</i>	<i>F</i>	<i>df</i>	<i>p_{ncor}</i>	ηp^2	<i>p_{cor}</i>
Global WM Capacity																				
Group	11.96	1	.001	.054	.002	26.48	1	.000	.114	.000	23.59	1	.000	.103	.000	10.78	1	.001	.049	.001
Genotype	0.47	2	.626	.004	.999	0.52	2	.593	.005	.999	0.13	1	.724	.001	.999	0.07	1	.791	.000	.791
Group*Genotype	4.87	2	.009	.044	.018	6.54	2	.002	.060	.006	2.30	1	.131	.011	.131	6.24	1	.013	.029	.026
Spatial WM (SWM)																				
Group	5.39	1	.021	.025	.042	13.52	1	.000	.062	.001	14.29	1	.000	.065	.000	3.33	1	.069	.016	.069
Genotype	2.99	2	.053	.028	.212	0.70	2	.498	.007	.999	0.02	1	.899	.000	.899	0.33	1	.566	.002	.999
Group*Genotype	3.18	2	.044	.029	.088	4.02	2	.019	.038	.057	3.23	1	.074	.016	.074	11.15	1	.001	.050	.004
Verbal WM (LNST)																				
Group	7.93	1	.005	.036	.005	17.55	1	.000	.079	.000	16.04	1	.000	.072	.000	8.17	1	.005	.037	.009
Genotype	0.88	2	.417	.008	.999	0.91	2	.403	.009	.999	0.23	1	.635	.001	.999	0.03	1	.859	.000	.859
Group*Genotype	3.26	2	.040	.030	.120	3.75	2	.025	.035	.100	1.72	1	.191	.008	.191	4.08	1	.045	.019	.090
Visuospatial WM (PAL)																				
Group	6.68	1	.010	.031	.010	12.85	1	.000	.056	.001	9.68	1	.002	.045	.006	7.88	1	.005	.036	.011
Genotype	0.32	2	.727	.003	.999	0.28	2	.760	.003	.999	0.10	1	.755	.000	.999	0.03	1	.854	.000	.854
Group*Genotype	2.19	2	.115	.020	.345	3.22	2	.042	.030	.168	0.01	1	.909	.000	.909	0.06	1	.803	.000	.999
Executive WM (Strategy)																				
Group	0.63	1	.428	.003	.428	5.81	1	.017	.028	.068	5.39	1	.021	.026	.063	1.84	1	.177	.009	.354
Genotype	0.86	2	.427	.008	.999	3.07	2	.048	.029	.192	0.09	1	.766	.000	.999	0.00	1	.964	.000	.964
Group*Genotype	5.55	2	.004	.050	.016	4.88	2	.009	.045	.027	0.66	1	.417	.003	.417	1.84	1	.176	.009	.352

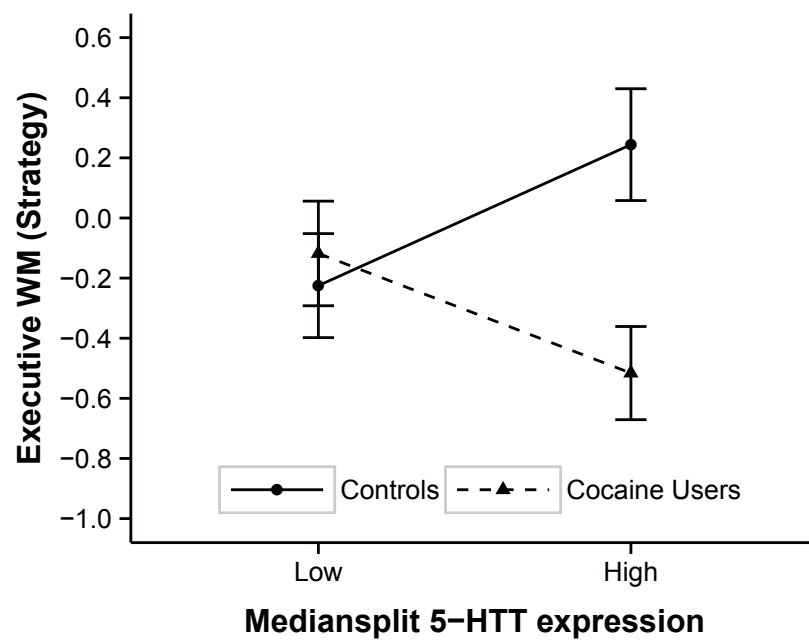
LNST: Letter-Number-Span-Task, PAL: Pair Associated Learning, *p_{ncor}*: uncorrected p-values, *p_{cor}*: p-values corrected for multiple comparisons, SWM: Spatial Working Memory, WM: working memory

Figure Legends

Fig. 1: Global working memory capacity scores for the three polymorphisms with significant *group*genotype* interactions (mean and *SE*, z-transformed and corrected for *age* and *years of education*). Significant genotype effects ($p < 0.05$) within each group are indicated with *, while trends for genotype effects ($p < 0.1$) are indicated with (*).

Fig. 2: Executive working memory function (Spatial Working Memory strategy scores) in cocaine users and controls stratified for high vs. low of 5-HTT mRNA expression (mean and *SE*, z-transformed and corrected for *age* and *years of education*).





Supplementary Information

Serotonin transporter and tryptophan hydroxylase gene variations mediate working memory deficit of cocaine users

Michael M Havranek, Matthias Vonmoos, Christian P Müller, Jessica R Büetiger, Eve Tasiudi, Lea M Hulka, Katrin H Preller, Rainald Mössner, Edna Grünblatt, Erich Seifritz, Boris B Quednow

Genotyping

DNA for the SNP genotyping was extracted either from EDTA anticoagulated blood samples or from immortalized lymphoblastoids cell cultures after transforming the lymphocytes with Epstein-Barr virus. The isolation of the DNA followed the QIAGEN protocol for the Blood & Cell Culture DNA Isolation Maxi Kit (QIAGEN, Hilden, Germany). To determine the 5-HTTLPR, PCR was performed in a final solution of 10µl, using 12.5ng of DNA, 2.5mM dNTPs, 0.5µM of each primer (5'-TCC TCC GCT TTG GCG CCT CTT CC-3' (forward primer); 5'-TGG GGG TTG CAG GGG AGA TCC TG-3' (reverse primer)), 0.5 U of Taq polymerase from Qiagen with addition of 2µl Q-solution. The initial denaturation step of five minutes at 95°C was followed by 40 cycles each of 95°C for 30 seconds, 65.5°C for 90 seconds and 72°C for 1 minute. PCR was terminated by a final extension step at 72°C for 10 minutes. 7.5 µl of the PCR product were utilized to separate the 512bp (long-) and 469 bp (short-) alleles of the 5-HTTLPR.

For the VNTR In2 polymorphisms, PCR was performed in a final solution of 10µl, using 12.5ng of DNA, 2.5mM dNTPs, 0.3µM of each primer (5'-GTC AGT ATC ACA GGC TGC GAG-3' (forward primer), 5'-TGT TCC TAG TCT TAC GCC AGT G-3' (reverse primer)) and 0.5 U of Taq-polymerase (Qiagen). The initial denaturation step of three minutes at 95°C was followed by 45 cycles each of 95°C for 30 seconds, 60°C for 30 seconds and 72°C for 45 seconds. PCR was terminated by a final extension step at 72°C for 10 minutes. We took 7.5 µl of the resulting PCR product and

separated the different alleles on a 2.5% LE agarose gel (Biozym, Hess. Oldendorf, Germany) by electrophoresis. The length of the alleles were: 251bp (9x), 268bp (10x) and 302bp (12x), respectively.

To determine the SNPs in the TPH2 gene (rs4570625, rs1386497), PCR was performed using 12.5ng of DNA and the SNP Genotyping Assays provided by Applied Biosystems (Foster City, CA, USA). The procedure followed the protocol for Taqman[®] SNP-Genotyping with the use of Taqman Universal PCR MasterMix and No AmpErase UNG. Each of the assays consisted of the unlabeled forward and reverse primers and two reporters that were dye-labeled with FAM[™] and VIC[®] and were designed for allelic discrimination of specific SNPs. Both alleles were scored in a single well by measuring the fluorescence at the end of the PCR using a Tecan Ultra 384 reader (Tecan, Crailsheim, Germany). Excitation- and emission-wavelengths were 485 and 535 nm for the FAM-labeled probes and 535 and 590 nm for the VIC-labeled probes, respectively.

Tab. S1: Substance use (means and *SD*).

Substance use	Controls (<i>n</i> =94)	Cocaine users (<i>n</i> =126)
Weekly nicotine use, cigarettes, mean (<i>SD</i>)	62.0 (66.5)	92.6 (75.4)
Weekly alcohol dose, g, mean (<i>SD</i>)	123.6 (135.9)	180.9 (194.8)
Weekly cocaine dose, g, mean (<i>SD</i>)	-	2.4 (5.1)
Years of cocaine use, mean (<i>SD</i>)	-	7.0 (5.4)
Cocaine hair analysis, pg/mg, mean (<i>SD</i>)	-	7804.9 (18713.9)
Cumulative cocaine dose, g, mean (<i>SD</i>)	-	2109.6 (5799.7)

Tab. S2: Genotype frequencies of all four investigated polymorphisms in total sample, in controls and in cocaine users (HWE = Hardy-Weinberg Equilibrium).

Polymorphism	Genotype frequencies %		Deviation from HWE?	in controls %	in users %	Association with group?
5-HTTLPR	L/L	33.2	$\chi^2(1) = .498, p = .481$	33.0	33.3	$\chi^2(1) = .00, p = .990$
	S/L	51.2		51.6	50.8	
	S/S	15.7		15.4	15.9	
VNTR In2	12/12	37.3	$\chi^2(1) = .707, p = .401$	35.2	38.8	$\chi^2(1) = .64, p = .423$
	12/9+10	44.3		44.0	44.6	
	9+10/9+10	18.4		20.9	16.5	
TPH2 (rs4570625)	G/G	62.4	$\chi^2(1) = 1.508, p = .219$	63.6	61.5	$\chi^2(1) = 1.07, p = .301$
	G/T	31.4		31.8	32.0	
	T/T	6.2		4.5	6.6	
TPH2 (rs1386497)	A/A	71.6	$\chi^2(1) = .584, p = .445$	68.5	75.0	$\chi^2(1) = .25, p = .619$
	A/C	24.8		28.3	22.6	
	C/C	3.6		3.3	2.4	

Tab. S3: Cocaine use across genotype groups (cocaine hair analysis, pg/mg, and cumulative cocaine dose in g, means and *SD*).

Cocaine use	Genotype	Cocaine hair analysis, pg/mg, mean (<i>SD</i>)	<i>F</i>	<i>df, df_{err}</i>	<i>p</i>
5-HTTLPR	L/L	8615 (17995)	0.280	2, 121	0.756
	S/L	6648 (17732)			
	S/S	9887 (23440)			
VNTR In2	12/12	2039 (5583)	0.354	2, 116	0.703
	12/9+10	6907 (15986)			
	9+10/9+10	9458 (24490)			
TPH2 (rs4570625)	G/G	6695 (21799)	0.901	1, 117	0.345
	G/T + T/T	10117 (21798)			
TPH2 (rs1386497)	A/A	7913 (17236)	0.000	1, 120	0.991
	A/C + C/C	7954 (22777)			
Cocaine use	Genotype	Cumulative cocaine dose, g, mean (<i>SD</i>)	<i>F</i>	<i>df, df_{err}</i>	<i>p</i>
5-HTTLPR	L/L	2631 (6030)	0.393	2, 122	0.676
	S/L	2028 (6220)			
	S/S	1226 (3461)			
VNTR In2	12/12	2039 (5583)	0.058	2, 117	0.944
	12/9+10	2374 (7036)			
	9+10/9+10	1941 (2590)			
TPH2 (rs4570625)	G/G	2560 (7024)	0.767	1, 118	0.383
	G/T + T/T	1591 (3534)			
TPH2 (rs1386497)	A/A	1946 (5216)	0.287	1, 121	0.593
	A/C + C/C	2586 (5844)			

Tab. S4: mRNA expression levels in controls and cocaine users and across genotype groups (means and *SD*).

mRNA expression	Genotype	Peripheral 5-HTT mRNA expression, mean (<i>SD</i>)			<i>F</i>	<i>df, df_{err}</i>	<i>p</i>
		in controls	in cocaine users				
5-HTTLPR	L/L	1.28 (.58)	1.21 (.41)	Factor group	0.011	1, 106	0.916
	S/L	1.23 (.59)	1.20 (.59)	Factor genotype	0.161	2, 106	0.852
	S/S	1.27 (.80)	1.34 (.68)				
VNTR In2	12/12	1.24 (.75)	1.38 (.70)	Factor group	0.86	1, 102	0.77
	12/9+10	1.21 (.46)	1.05 (.40)	Factor genotype	1.165	2, 102	0.316
	9+10/9+10	1.30 (.67)	1.22 (.40)				
TPH2 (rs4570625)	G/G	1.17 (.60)	1.22 (.55)	Factor group	1.518	1, 105	0.221
	G/T + T/T	1.60 (.58)	1.24 (.44)	Factor genotype	3.389	1, 105	0.068
TPH2 (rs1386497)	A/A	1.32 (.49)	1.21 (.51)	Factor group	0.258	1, 105	0.612
	A/C + C/C	1.21 (.75)	1.21 (.55)	Factor genotype	0.214	1, 105	0.645

Figure S1

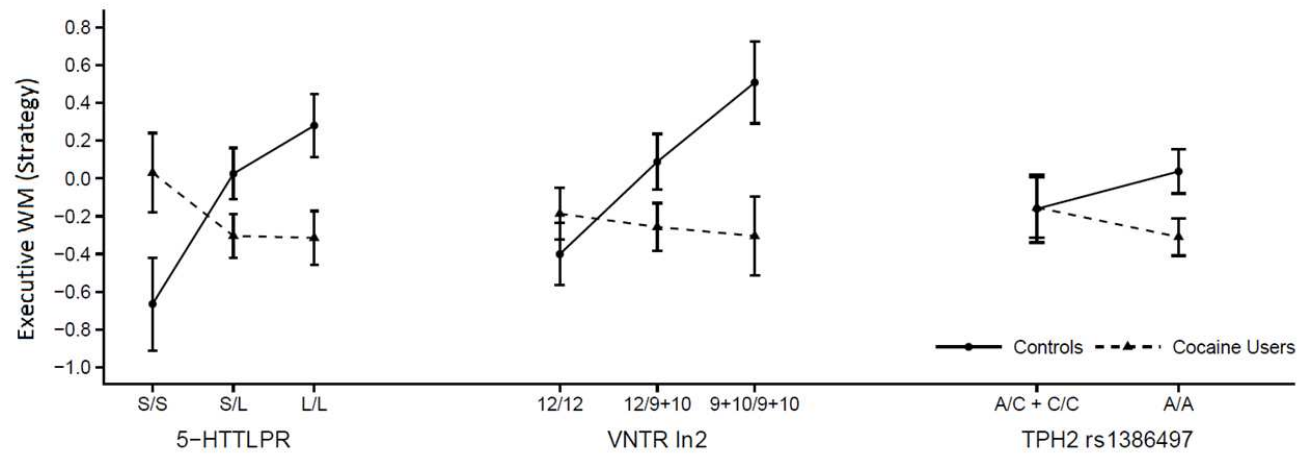
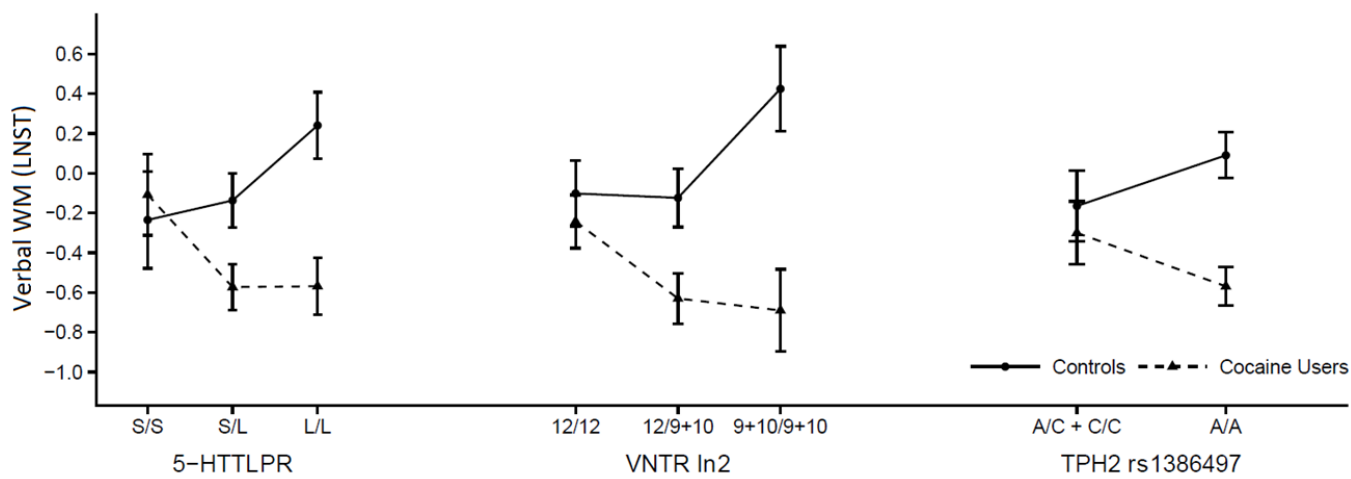
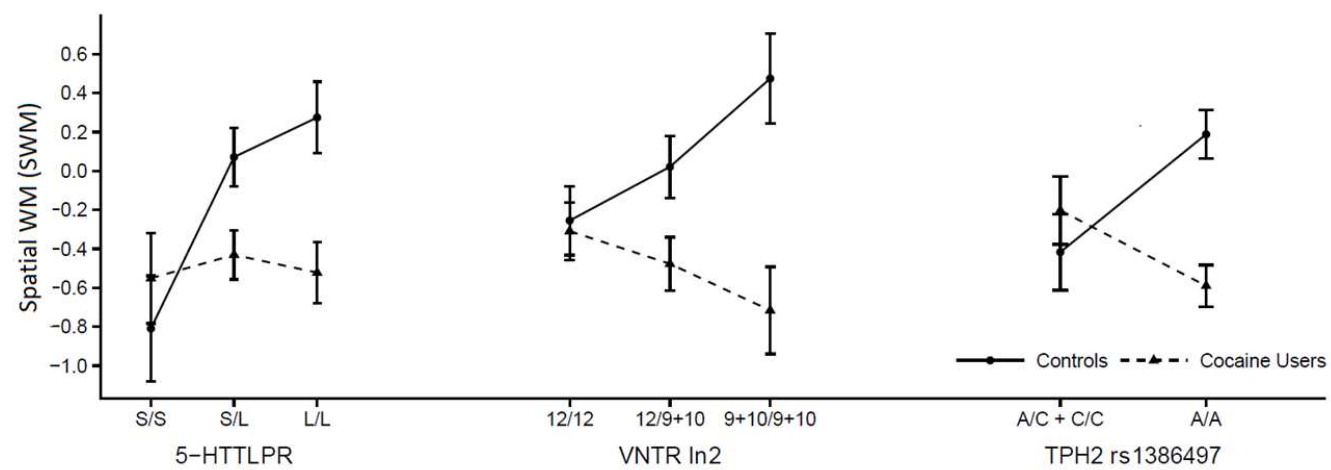


Fig. S1: Executive WM scores for the three polymorphisms with significant *group*genotype* interactions in 5-HTTLPR and VNTR In2 but not in TPH2 rs1386497 (mean and SE, z-transformed and corrected for age and years of education).

Figure S2



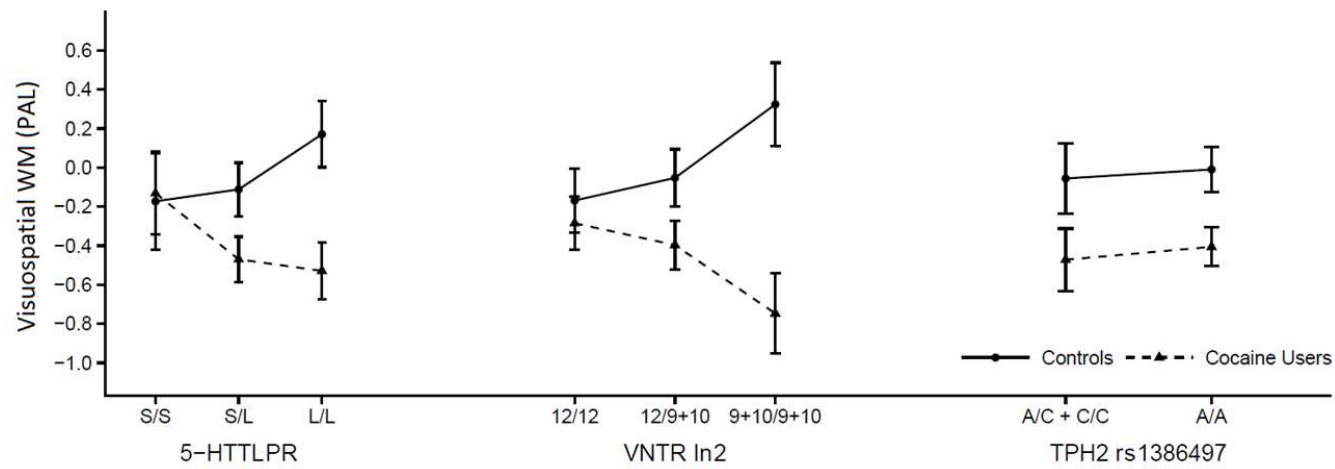


Fig. S2: Spatial WM (A), verbal WM (B) and visuospatial WM scores (C) for 5-HTTLPR, VNTR In2 and TPH2 rs1386497 (mean and *SE*, *z*-transformed and corrected for *age* and *years of education*).

Figure S3

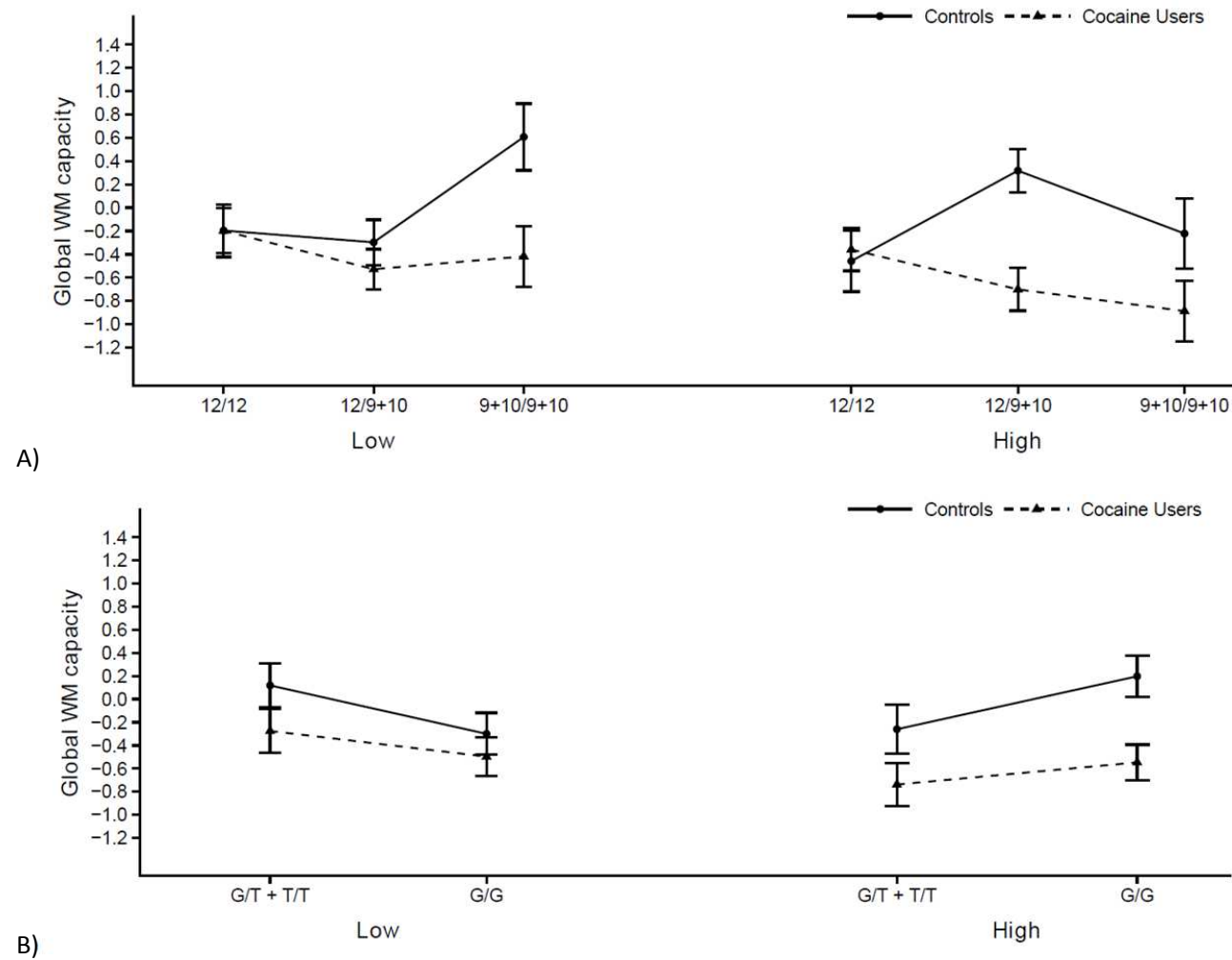


Fig. S3: Global WM scores for the mediansplit (low vs. high) of 5-HTT mRNA expression in A) VNTR In2 and B) TPH2 rs4570625 (mean and SE, z-transformed and corrected for *age* and *years of education*).

Figure S4

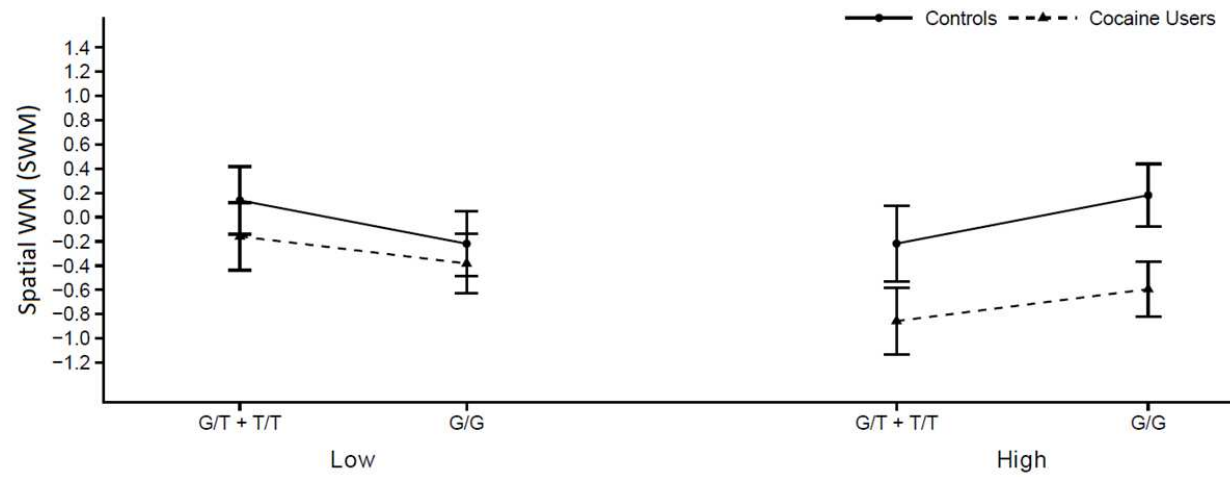
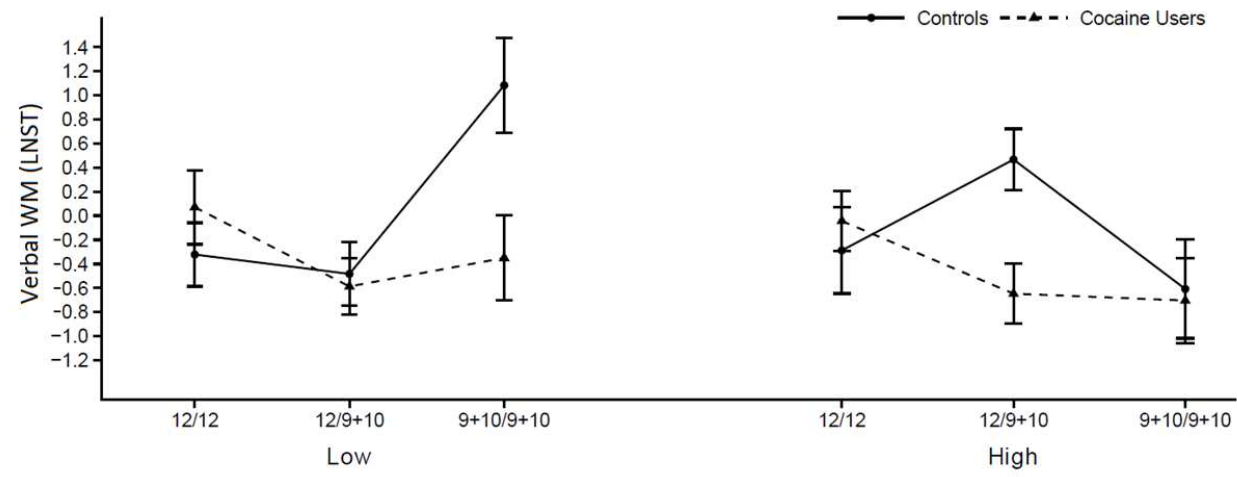
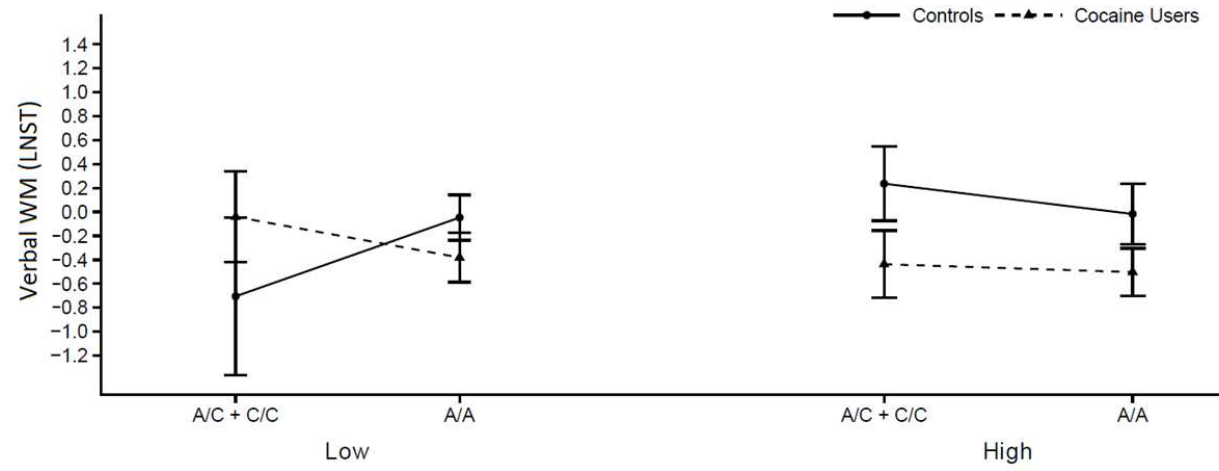


Fig. S4: Spatial WM scores for the mediansplit (low vs. high) of 5-HTT mRNA expression in TPH2 rs4570625 (mean and *SE*, z-transformed and corrected for *age* and *years of education*).

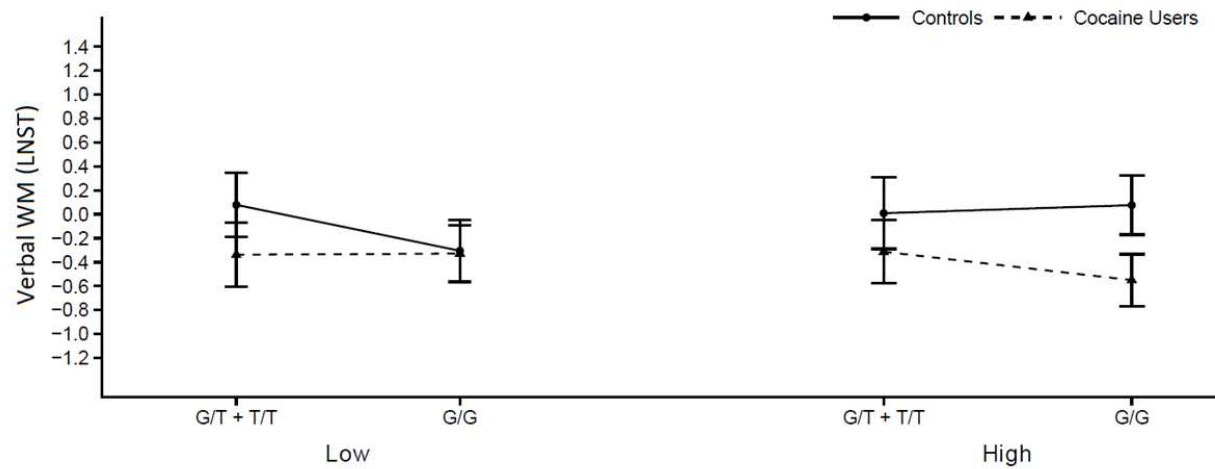
Figure S5



A)



B)



C)

Fig. S5: Verbal WM scores for the mediansplit (low vs. high) of 5-HTT mRNA expression in A) VNTR In2, B) TPH2 rs1386497 and C) TPH2 rs4570625 (mean and *SE*, z-transformed and corrected for *age* and *years of education*).

Figure S6

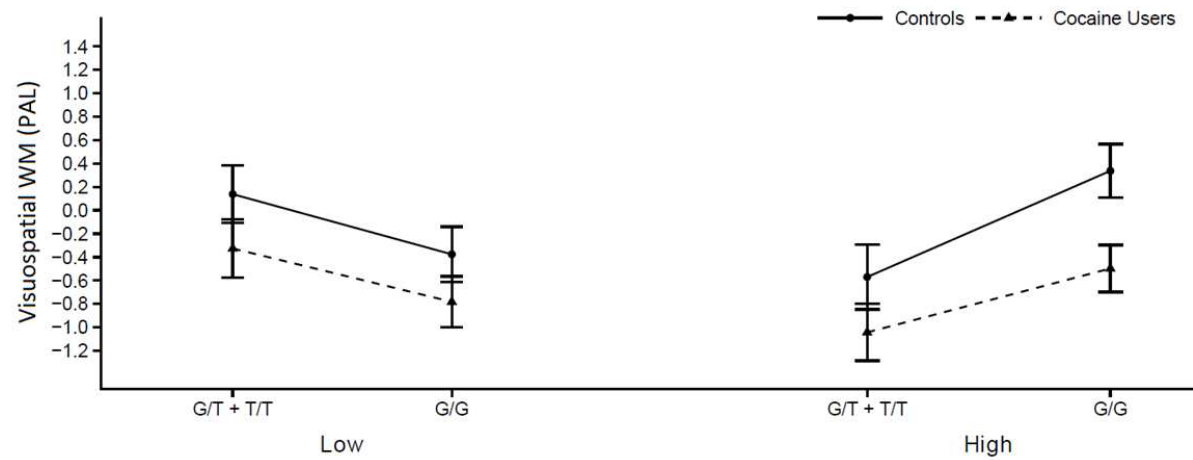


Fig. S6: Visuospatial WM scores for the mediansplit (low vs. high) of 5-HTT mRNA expression inTPH2 rs4570625 (mean and *SE*, z-transformed and corrected for *age* and *years of education*).

Figure S7

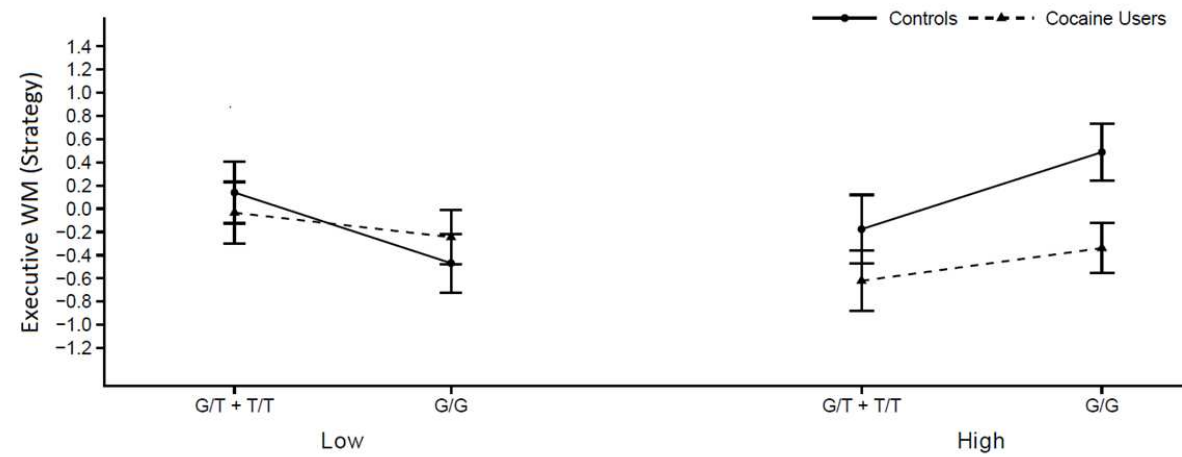


Fig. S7: Executive WM scores for the mediansplit (low vs. high) of 5-HTT mRNA expression inTPH2 rs4570625 (mean and *SE*, z-transformed and corrected for *age* and *years of education*).